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## Claims

- 1. A method for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said method comprising the steps of:
  - (a) providing a cell having:
- (i) a reporter gene operably linked to a DNA-binding-protein recognition site;
  - (ii) a first fusion gene capable of expressing a first fusion protein, said first fusion protein comprising a polypeptide fragment of Smad2 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site; and
  - (iii) a second fusion gene capable of expressing a second fusion protein, said second fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a gene activating moiety;
    - (b) exposing said cell to said compound; and
- (c) measuring reporter gene expression in said cell, a change in said reporter gene expression indicating said compound is capable of modulating TGFβ superfamily signalling.
- 2. A method for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said method comprising the steps of:
  - (a) providing a cell having:
- (i) a reporter gene operably linked to a DNA-binding-protein recognition site;
  - (ii) a first fusion gene capable of expressing a first fusion

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protein, said first fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site;

- (iii) a second fusion gene capable of expressing a second fusion protein, said second fusion protein comprising a polypeptide fragment of Smad2 covalently bonded to a gene activating moiety;
  - (b) exposing said cell to said compound; and
- (c) measuring reporter gene expression in said cell, a change in said reporter gene expression indicating said compound is capable of modulating TGF-β superfamily signalling.
- 3. A method for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said method comprising the steps of:
  - (a) providing a cell having:
- (i) a reporter gene operably linked to a DNA-binding-protein recognition site;
- (ii) a first fusion gene capable of expressing a first fusion protein, said first fusion protein comprising a polypeptide fragment of Smad3 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site; and
- (iii) a second fusion gene capable of expressing a second fusion protein, said second fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a gene activating moiety;
  - (b) exposing said cell to said compound; and
  - (c) measuring reporter gene expression in said cell, a change in said

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reporter gene expression indicating said compound is capable of modulating TGF-  $\beta$  superfamily signalling.

- 4. A method for detecting a compound capable of modulating TGF-β superfamily signalling, said method comprising the steps of:
  - (a) providing a cell having:
- (i) a reporter gene operably linked to a DNA-binding-protein recognition site;
- (ii) a first fusion gene capable of expressing a first fusion protein, said first fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site;
- (iii) a second fusion gene capable of expressing a second fusion protein, said second fusion protein comprising a polypeptide fragment of Smad3 covalently bonded to a gene activating moiety;
  - (b) exposing said cell to said compound; and
- (c) measuring reporter gene expression in said cell, a change in said reporter gene expression indicating said compound is capable of modulating TGF-β superfamily signalling.
- 5. A cell for detecting a compound capable of modulating TGF-β superfamily signalling, said cell having:
- (a) a reporter gene operably linked to a DNA-binding-protein recognition site;
  - (b) a first fusion gene capable of expressing a first fusion protein, said

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first fusion protein comprising a polypeptide fragment of Smad2 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site; and

- (c) a second fusion gene capable of expressing a second fusion protein, said second fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a gene activating moiety.
- 6. A cell for detecting a compound capable of modulating TGF-β superfamily signalling, said cell having:
- (a) a reporter gene operably linked to a DNA-binding-protein recognition site;
- (b) a first fusion gene capable of expressing a first fusion protein, said first fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site; and
- (c) a second fusion gene capable of expressing a second fusion protein, said second fusion protein comprising a polypeptide fragment of Smad2 covalently bonded to a gene activating moiety.
- 7. A cell for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said cell having:
- (a) a reporter gene operably linked to a DNA-binding-protein recognition site;
- (b) a first fusion gene capable of expressing a first fusion protein, said first fusion protein comprising a polypeptide fragment of Smad3 covalently

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bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site; and

- (c) a second fusion gene capable of expressing a second fusion protein, said second fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a gene activating moiety.
- 8. A cell for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said cell having:
- (a) a reporter gene operably linked to a DNA-binding-protein recognition site;
- (b) a first fusion gene capable of expressing a first fusion protein, said first fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site; and
- (c) a second fusion gene capable of expressing a second fusion protein, said second fusion protein comprising a polypeptide fragment of Smad3 covalently bonded to a gene activating moiety.
- 9. A method for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said method comprising the steps of:
- (a) providing a first polypeptide, said first polypeptide comprising a polypeptide fragment of FAST-1;
- (b) providing a second polypeptide, said second polypeptide comprising a polypeptide fragment of Smad2;
  - (c) exposing said first polypeptide to said second polypeptide and to said

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compound; and

- (d) measuring the level of interaction between said first polypeptide and said second polypeptide, an alteration in said level of interaction indicating said compound is capable of modulating TGF-β superfamily signalling.
- 10. A method for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said method comprising the steps of:
- (a) providing a first polypeptide, said first polypeptide comprising a polypeptide fragment of Smad2;
- (b) providing a second polypeptide, said second polypeptide comprising a polypeptide fragment of FAST-1;
- (c) exposing said first polypeptide to said second polypeptide and to said compound; and
- (d) measuring the level of interaction between said first polypeptide and said second polypeptide, an alteration in said level of interaction indicating said compound is capable of modulating TGF-β superfamily signalling.
- 11. A method for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said method comprising the steps of:
- (a) providing a first polypeptide, said first polypeptide comprising a polypeptide fragment of FAST-1;
- (b) providing a second polypeptide, said second polypeptide comprising a polypeptide fragment of Smad3;
- (c) exposing said first polypeptide to said second polypeptide and to said compound; and

- (d) measuring the level of interaction between said first polypeptide and said second polypeptide, an alteration in said level of interaction indicating said compound is capable of modulating TGF- $\beta$  superfamily signalling.
- 12. A method for detecting a compound capable of modulating TGF-β superfamily signalling, said method comprising the steps of:
- (a) providing a first polypeptide, said first polypeptide comprising a polypeptide fragment of Smad3;
- (b) providing a second polypeptide, said second polypeptide comprising a polypeptide fragment of FAST-1;
- (c) exposing said first polypeptide to said second polypeptide and to said compound; and
- (d) measuring the level of interaction between said first polypeptide and said second polypeptide, an alteration in said level of interaction indicating said compound is capable of modulating TGF-β superfamily signalling.
- 13. A method for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said method comprising the steps of:
- (a) providing a reporter gene operably linked to a DNA-binding-protein recognition site;
- (b) providing a first fusion protein, said first fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site;
  - (c) providing a second fusion protein, said second fusion protein

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comprising a polypeptide fragment of Smad2 covalently bonded to a gene activating moiety;

- (d) exposing said first fusion protein to said second fusion protein, to said reporter gene, and to said compound; and
- (e) measuring the reporter gene expression, a change in said reporter gene expression indicating a compound capable of modulating TGF- $\beta$  superfamily signalling.
- 14. A method for detecting a compound capable of modulating TGF-β superfamily signalling, said method comprising the steps of:
- (a) providing a reporter gene operably linked to a DNA-binding-protein recognition site;
- (b) providing a first fusion protein, said first fusion protein comprising a polypeptide fragment of Smad2 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site;
- (c) providing a second fusion protein, said second fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a gene activating moiety;
- (d) exposing said first fusion protein to said second fusion protein, to said reporter gene, and to said compound; and
- (e) measuring the reporter gene expression, a change in said reporter gene expression indicating a compound capable of modulating TGF- $\beta$  superfamily signalling.

- 15. A method for detecting a compound capable of modulating TGF- $\beta$  superfamily signalling, said method comprising the steps of:
- (a) providing a reporter gene operably linked to a DNA-binding-protein recognition site;
- (b) providing a first fusion protein, said first fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site;
- (c) providing a second fusion protein, said second fusion protein comprising a polypeptide fragment of Smad3 covalently bonded to a gene activating moiety;
- (d) exposing said first fusion protein to said second fusion protein, to said reporter gene, and to said compound; and
- (e) measuring the reporter gene expression, a change in said reporter gene expression indicating a compound capable of modulating TGF- $\beta$  superfamily signalling.
- 16. A method for detecting a compound capable of modulating TGF-β superfamily signalling, said method comprising the steps of:
- (a) providing a reporter gene operably linked to a DNA-binding-protein recognition site;
- (b) providing a first fusion protein, said first fusion protein comprising a polypeptide fragment of Smad3 covalently bonded to a binding moiety, said binding moiety capable of specifically binding to said DNA-binding-protein recognition site;

- (c) providing a second fusion protein, said second fusion protein comprising a polypeptide fragment of FAST-1 covalently bonded to a gene activating moiety;
- (d) exposing said first fusion protein to said second fusion protein, to said reporter gene, and to said compound; and
- (e) measuring the reporter gene expression, a change in said reporter gene expression indicating a compound capable of modulating TGF- $\beta$  superfamily signalling.
- 17. A method for diagnosing a mammal having or likely to develop a disorder involving abnormal TGF-β superfamily signalling, said method comprising determining whether said mammal has a mutation in a gene encoding FAST-1.
- 18. A method for diagnosing a mammal having or likely to develop a disorder involving abnormal TGF- $\beta$  superfamily signalling, said method comprising determining whether said mammal has an altered level of expression of FAST-1.
- 19. A substantially pure FAST-1 protein or polypeptide fragment thereof, wherein said protein or said polypeptide fragment is from a mammal, wherein said protein or polypeptide fragment is for use in modulating TGF- $\beta$  superfamily signalling.
- 20. A substantially pure polypeptide fragment, wherein said polypeptide fragment is a polypeptide fragment of FAST-1, wherein said FAST-1

is from *Xenopus*, wherein said polypeptide fragment comprises the Smad Interaction Domain (SID), wherein said polypeptide fragment is for use in modulating TGF- $\beta$  superfamily signalling.

- 21. A substantially pure polypeptide, wherein said polypeptide has
  5 about 50% or greater amino acid sequence identity to the amino acid sequence of a substantially pure mammalian FAST-1 protein, or polypeptide fragment thereof, wherein said protein or said polypeptide fragment is for use in modulating TGF-β superfamily signalling.
  - 22. A substantially pure polypeptide, wherein said polypeptide has about 75% or greater amino acid sequence identity to the amino acid sequence of a substantially pure mammalian FAST-1 protein, or polypeptide fragment thereof, wherein said protein or said polypeptide fragment is for use in modulating TGF- $\beta$  superfamily signalling.
  - 23. A substantially pure polypeptide, wherein said polypeptide has about 90% or greater amino acid sequence identity to the amino acid sequence of a substantially pure mammalian FAST-1 protein, or polypeptide fragment thereof, wherein said protein or said polypeptide fragment is for use in modulating TGF- $\beta$  superfamily signalling.
- 24. A substantially pure nucleic acid, wherein said nucleic acid encodes
   20 a mammalian FAST-1 protein, or polypeptide fragment thereof, wherein said
   protein or said polypeptide fragment is for use in modulating TGF-β superfamily

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- 25. A vector comprising a substantially pure nucleic acid, wherein said nucleic acid encodes a mammalian FAST-1 protein, or polypeptide fragment thereof, wherein said protein or said polypeptide fragment is for use in modulating TGF- $\beta$  superfamily signalling, and wherein said vector is capable of directing expression of said protein or said polypeptide fragment in a cell containing said vector.
- 26. A vector comprising a substantially pure nucleic acid, wherein said nucleic acid encodes a FAST-1 Smad Interaction Domain (SID), wherein said SID is for use in modulating TGF- $\beta$  superfamily signalling, and wherein said vector is capable of directing expression of said SID in a cell containing said vector.
- 27. A cell that contains a vector comprising a substantially pure nucleic acid, wherein said nucleic acid encodes a mammalian FAST-1 protein, or polypeptide fragment thereof, wherein said protein or said polypeptide fragment is for use in modulating TGF-β superfamily signalling, and wherein said vector is capable of directing expression of said protein or said polypeptide fragment.
- 28. A method of modulating TGF- $\beta$  superfamily signalling in a cell, said method comprising providing a cell with a substantially pure FAST-1 protein, or polypeptide fragment thereof, wherein said FAST-1 protein or polypeptide fragment is provided intracellularly, and wherein said FAST-1 protein or polypeptide fragment is sufficient to modulate TGF- $\beta$  superfamily signalling in a

29. A method of modulating TGF- $\beta$  superfamily signalling in a cell, said method comprising introducing, into a cell, a vector comprising a substantially pure nucleic acid, wherein said nucleic acid encodes a substantially pure FAST-1 protein, or polypeptide fragment thereof, wherein said vector is capable of directing expression of said protein or said polypeptide fragment in a cell containing said vector, and wherein expression of said FAST-1 protein or polypeptide fragment is sufficient to modulate TGF- $\beta$  superfamily signalling in a cell.